

PCT

RAW SEQUENCE LISTING DATE: 07/16/2004
PATENT APPLICATION: US/10/501,196 TIME: 16:55:45

Input Set : A:\NEB-198-PJP.txt

Output Set: N:\CRF4\07162004\J501196.raw

```
3 <110> APPLICANT: NEW ENGLAND BIOLABS, INC.
             SAMUELSON, JAMES
             XU, SHUANG-YONG
      7 <120> TITLE OF INVENTION: ALTERATION OF RESTRICTION ENDONUCLEASE SPECIFICITY BY
GENETIC SELECTION
     9 <130> FILE REFERENCE: NEB-198-PCT
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/501,196
C--> 12 <141> CURRENT FILING DATE: 2004-07-09
     14 <150> PRIOR APPLICATION NUMBER: US 60/347,403
     15 <151> PRIOR FILING DATE: 2002-01-10
     17 <160> NUMBER OF SEQ ID NOS: 2
     19 <170> SOFTWARE: PatentIn version 3.1
     21 <210> SEQ ID NO: 1
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     22 <211> LENGTH: 612
     23 <212> TYPE: DNA
     24 <213> ORGANISM: Bacillus stearothermophilus Y406
     26 <220> FEATURE:
     27 <221> NAME/KEY: CDS
     28 <222> LOCATION: (1)..(612)
     29 <223> OTHER INFORMATION:
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     34 Met Arg Ile Val Glu Val Tyr Ser His Leu Asn Gly Leu Glu Tyr Ile
                                                                               96
     37 caa gtt cac ttg cca cat att tgg gaa gaa att caa gaa att att gtt
     38 Gln Val His Leu Pro His Ile Trp Glu Glu Ile Gln Glu Ile Ile Val
                                        25
     41 tct att gac gca gaa gct tgt aga acg aag gaa tca aaa gaa aag aca
                                                                              144
     42 Ser Ile Asp Ala Glu Ala Cys Arg Thr Lys Glu Ser Lys Glu Lys Thr
                                                                              192
     45 aaa caa gga caa ata ctt tat agt ccc gta gct tta aat gaa gca ttc
     46 Lys Gln Gly Gln Ile Leu Tyr Ser Pro Val Ala Leu Asn Glu Ala Phe
     47
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     49 aag gaa aaa tta gaa gca aaa ggt tgg aaa gaa agt cga aca aac tat
                                                                              240
     50 Lys Glu Lys Leu Glu Ala Lys Gly Trp Lys Glu Ser Arg Thr Asn Tyr
                                                75
                                                                              288
     53 tat gtg, act gct gac cca aag ctg att cgt gaa aca tta tca ctt gaa
     54 Tyr Val Thr Ala Asp Pro Lys Leu Ile Arg Glu Thr Leu Ser Leu Glu
                                            90
     57 cca gag gaa caa aag aaa gtg att gaa gcc gca gga aaa gaa gca tta
                                                                              336
     58 Pro Glu Glu Gln Lys Lys Val Ile Glu Ala Ala Gly Lys Glu Ala Leu
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     61 aag tot tat aat caa acg gat ttt gta aaa gat aga gtg gca ata gaa
                                                                              384
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62 Lys Ser Tyr Asn Gln Thr Asp Phe Val Lys Asp Arg Val Ala Ile Glu

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65 gtt caa ttc gga aaa tat tct ttt gtc gct tat gac ctt ttc gtc aaa	432
66 Val Gln Phe Gly Lys Tyr Ser Phe Val Ala Tyr Asp Leu Phe Val Lys	
67 130 135 140	
69 cac atg gct ttc tat gtt agt gat aaa att gac gtt ggt gtc gaa ata	480
70 His Met Ala Phe Tyr Val Ser Asp Lys Ile Asp Val Gly Val Glu Ile	
71 145 150 155 160	
73 ttg cca atg aag gaa tta tca aaa gaa atg tct tcg gga atc agt tat	528
74 Leu Pro Met Lys Glu Leu Ser Lys Glu Met Ser Ser Gly Ile Ser Tyr	
75 165 170 175	
77 tac gaa ggt gaa tta tac aat gtg ata cgg caa ggt cgt ggc gtt cct	576
78 Tyr Glu Gly Glu Leu Tyr Asn Val Ile Arg Gln Gly Arg Gly Val Pro	
73	612
81 gcc gtt ccg ttg gtt tta atc ggg att gcc cct taa	012
82 Ala Val Pro Leu Val Leu Ile Gly Ile Ala Pro	
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86 <210> SEQ ID NO: 2	
87 <211> LENGTH: 203	
88 <212> TYPE: PRT	
89 <213> ORGANISM: Bacillus stearothermophilus Y406	
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98 20 25 30	
101 Ser Ile Asp Ala Glu Ala Cys Arg Thr Lys Glu Ser Lys Glu Lys Thr	
102 35 40 45	
105 Lys Gln Gly Gln Ile Leu Tyr Ser Pro Val Ala Leu Asn Glu Ala Phe	
106 50 55 60	
109 Lys Glu Lys Leu Glu Ala Lys Gly Trp Lys Glu Ser Arg Thr Asn Tyr	
110 65 70 75 80	
113 Tyr Val Thr Ala Asp Pro Lys Leu Ile Arg Glu Thr Leu Ser Leu Glu	
113 Tyl Val Till Ala Asp F10 Hys hed T1c Alg C1d Till Zed Sel Zed C1d  114 85 90 95	
	•
117 Pro Glu Glu Gln Lys Lys Val Ile Glu Ala Ala Gly Lys Glu Ala Leu	
117 Pro Glu Glu Gln Lys Lys Val Ile Glu Ala Ala Gly Lys Glu Ala Leu 118 100 105 110	
117 Pro Glu Glu Gln Lys Lys Val Ile Glu Ala Ala Gly Lys Glu Ala Leu 118 100 105 110 121 Lys Ser Tyr Asn Gln Thr Asp Phe Val Lys Asp Arg Val Ala Ile Glu	
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VERIFICATION SUMMARY

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L:11 M:270 C: Current Application Number differs, Replaced Current Application Number

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:32 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:29